



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ISSUE 22
AMDIS: Setting Up and Running a Deconvolution and Target Analysis – Part 3
In the first two parts of the discussion of setting up AMDIS the basics of running an analysis with retention index filtering as well as a simple analysis were discussed. In this final part of the set-up section, instrument settings and filtering will be covered.
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ISSUE 21
AMDIS: Setting Up and Running a Deconvolution and Target Analysis – Part 2
In the previous part of this AMDIS discussion, the setting up and running of a simple analysis was discussed. However, to take full advantage of AMDIS it is far better to make use not only of the mass spectral information, but also the retention time data. In Part 2 of the "Setting Up and Running a Deconvolution and Target Analysis", we will discuss setting up and using the retention information and briefly discuss the various parameters that can be set by the user.
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ISSUE 20
AMDIS: Setting Up and Running a Deconvolution and Target Analysis – Part 1
AMDIS is designed to do analysis for target compounds in a GC/MS data file. To do the analysis you need a completed data file and a target library. You can put a target library together from standards using AMDIS or by extracting spectra from another library such as NIST11. The download package contains a small library that can be used for the data files in the tutorial with the download package. The examples used here will come from the tutorial package so that if you choose you may download the entire package and run the data for yourself.
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
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